

CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/844,684

**ENTERED**

CRF Processing Date: 1/31/2002  
 Edited by: \_\_\_\_\_  
 Verified by: ME (STIC staff)

01/E 0570 #8  
028

☐ Changed a file from non-ASCII to ASCII

☒ Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_

☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_

☐ Added the mandatory heading and subheadings for "Current Application Data".

☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_

☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_

☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_

☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_

☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_

☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_

☐ Inserted mandatory headings, specifically: \_\_\_\_\_

☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_

☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_

☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_

☐ Other: \_\_\_\_\_



OIEP

## RAW SEQUENCE LISTING

DATE: 01/31/2002

PATENT APPLICATION: US/09/844,684

TIME: 20:32:00

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\01312002\I844684.raw

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5 <110> APPLICANT: GEMINI SCIENCE, INC.
6     LA JOLLA INSTITUTE FOR ALLERGY AND IMMUNOLOGY
8 <120> TITLE OF INVENTION: HUMAN ANTI-CD40 ANTIBODIES AND METHODS OF MAKING SAME
10 <130> FILE REFERENCE: 21286/0276339
12 <140> CURRENT APPLICATION NUMBER: US 09/844,684
13 <141> CURRENT FILING DATE: 2001-04-27
15 <150> PRIOR APPLICATION NUMBER: US 60/200,601
16 <151> PRIOR FILING DATE: 2000-04-28
18 <160> NUMBER OF SEQ ID NOS: 15
20 <170> SOFTWARE: PatentIn Ver. 2.1
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24 <212> TYPE: DNA
25 <213> ORGANISM: Artificial Sequence
27 <220> FEATURE:
28 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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34 <210> SEQ ID NO: 2
35 <211> LENGTH: 41
36 <212> TYPE: DNA
37 <213> ORGANISM: Artificial Sequence
39 <220> FEATURE:
40 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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47 <211> LENGTH: 26
48 <212> TYPE: DNA
49 <213> ORGANISM: Artificial Sequence
51 <220> FEATURE:
52 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
54 <400> SEQUENCE: 3
55 gtgcacgccg ctggtcaggg cgcctg 26
58 <210> SEQ ID NO: 4
59 <211> LENGTH: 26
60 <212> TYPE: DNA
61 <213> ORGANISM: Artificial Sequence
W--> 62 <220> FEATURE:
63 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
65 <400> SEQUENCE: 4
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69 <210> SEQ ID NO: 5

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94 <211> LENGTH: 24
95 <212> TYPE: DNA
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98 <220> FEATURE:
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108 <213> ORGANISM: Artificial Sequence
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111 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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121 <213> ORGANISM: Artificial Sequence
123 <220> FEATURE:
124 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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130 <210> SEQ ID NO: 10
131 <211> LENGTH: 520
132 <212> TYPE: DNA
133 <213> ORGANISM: Homo sapiens
135 <400> SEQUENCE: 10
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138 agttcagcct ggggggtccc tgagactctc ctgtgcagtc tctggattca ccttcagtac 180
139 ctactggatg cactgggtcc gccaaagctcc aggggaagggg ctggtgtggg tctcacgtat 240
140 taatagtgat gggagtagca caacctacgc ggactccgtg aagggccgat tcaccatctc 300

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141 cagagacaac gccagaaca cgctgtatct gcaaatgaac agtctgagag ccgaggacac 360
142 ggctgtgtat tactgtgcaa gagatagagt actatggatc ggggagttat cctactacgg 420
143 tatggacgtc tggggccaag ggaccacggt caccgtctcc tcagctagca ccaagggccc 480
144 atcggctcttc cccctggcac cctcctccaa gagcacctct 520
147 <210> SEQ ID NO: 11
148 <211> LENGTH: 698
149 <212> TYPE: DNA
150 <213> ORGANISM: Homo sapiens
152 <400> SEQUENCE: 11
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155 caccctgtct gcatctgtag gagacagagt caccatcact tgccgggcca gtcagagtat 180
156 tagtaactgg ttggcctggg atcagcagaa accagggaag gccctaagc tctgtctcta 240
157 taaggcatct ggttttagaaa gtggggtccc atcaaggttc agcggcagtg gatctgggac 300
158 agaattcact ctcaccatca acagcctgca gcctgatgat ttgcaactt attactgcca 360
159 acagtctaata agttattcgt ggacgttcgg ccacgggacc aagggtgaaa tcaaacgtac 420
160 ggtggctgca ccatctgtct tcactctccc gccatctgat gagcagttga aatctggaac 480
161 tgctctgtgt gtgtgcctgc tgaataactt ctatcccaga gagggccaaag tacagtggaa 540
162 ggtggataac gccctccaat cgggtaactc ccaggagagt gtcacagagc aggacagcaa 600
163 ggacagcacc tacagcctca gcagcacctt gacgctgagc aaagcagact acgagaaaca 660
164 caaagtctac gcctgcgaag tcacccatca gggcctga 698
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168 <211> LENGTH: 580
169 <212> TYPE: DNA
170 <213> ORGANISM: Homo sapiens
172 <400> SEQUENCE: 12
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174 tgtggctatt ttaaaagggtg tccagtgtga ggtgcagctg ttggagtctg ggggaggctt 120
175 ggtacagcct ggggggtccc tgagactctc ctgtgcagcc tctggattcg cttttagcag 180
176 ctatgccatg agctgggtcc gccaggctcc aggggaagggg ctggagtggg tctcagctat 240
177 tagtggtagt ggtggtagca catactacgc agactccgtg aagggccggt tcaccatctc 300
178 cagagacaat tccaagaaca cgctgtatct gcaaatgaac agcctgagag ccgaggacac 360
179 ggccgtatat tactgtgcga aagatggggg gtactatggt tcggggagtt atgggtactt 420
180 tgactactgg ggccaggga cctggtcac cgtctcctca gctagcacca agggcccatc 480
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188 <212> TYPE: DNA
189 <213> ORGANISM: Homo sapiens
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194 gatgaccag tctccatctt ccgtgtctgc atctgcagga gacagagtca ccatcacttg 180
195 tcgggcgagt cagggtatta gcagctgggt agcctgggtat caacagaaac cagggaaaag 240
196 ccctaagctc ctgatctatg ctggatccag ttgcaaaagt ggggtcccat caaggttcag 300
197 cggcagtgga ttggggacag atttcaactc caccatcggc agcctgcagc ctgaagattt 360
198 tgcaacttac tattgtcaac aggtagcag ttccctcgg acgttcggcc aaggggacca 420
199 ggtggagatc aaacgtacgg tggtgcacc atctgtcttc atcttccgc catctgatga 480

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200 gcagttgaaa tctggaactg cctctgttgt gtgcctgctg aataacttct atcccagaga 540
201 ggccaaagta cagtgggaagg tggataacgc cctccaatcg ggtaactccc aggagagtgt 600
202 cacagagcag gacagcaagg acagcaccta cagcctcagc agcacccctga cgctgagcaa 660
203 agcagactac gagaaacaca aagtctacgc ctgcgaagtc acccatcagg gcctga 716
206 <210> SEQ ID NO: 14
207 <211> LENGTH: 630
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209 <213> ORGANISM: Homo sapiens
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213 ccgctcgacgg tgatcaggac tgaacagaga gaactcacca tggagtttgg gctgagctgg 120
214 ctttttcttg tggctatttt aaaagggtgc cagtgtgagg tgcagctgtt ggagtctggg 180
215 ggaggcttgg tacagcctgg ggggtccctg agactctcct gtgcagcctc tggattcacc 240
216 tttagcagct atgccatgag ctgggtccgc caggctccag ggaaggggct ggagtgggtc 300
217 tcagctatta gtggtagtgg tggtagcaca tactacgcag actccgtgaa yyyccggttc 360
218 accatctcca gagacaattc caagaacacg ctgtatctgc aaatgaacag cctgagagcc 420
219 gaggacacgg ccgtatatta ctgtgcgaaa gatggggggg actatgggtc ggggagttat 480
220 ggggtactttg actactgggg ccagggaacc ctggtcaccg tctcctcagc tagcaccaag 540
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222 ctgggctgcc tgggtcaagga ctacttcccc 630
225 <210> SEQ ID NO: 15
226 <211> LENGTH: 728
227 <212> TYPE: DNA
228 <213> ORGANISM: Homo sapiens
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233 atgcgacatc cagatgaccc agtctccatc ttccgtgtct ggatctgtag gagacagagt 180
234 caccatcact tgtcggggcg gtcagggtat tagcagctgg ttagcctggg atcagcagaa 240
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236 atcaagggtc agcggcagtg gatttgggac agatttcaact ctcaccatca gcagcctgca 360
237 gcctgaagat tttgcaactt actattgtca acaggctagc agtttccctc ggacattcgg 420
238 ccaagggacc aagggtggaga tcaaacgtac ggtggctgca ccactctgtc tcatcttccc 480
239 gccatctgat gagcagttga aatctggaac tgctctgtt gtgtgcctgc tgaataactt 540
240 ctatcccaga gaggccaaag tacagtggaa ggtggataac gccctccaat cgggtaactc 600
241 ccaggagagt gtcacagagc aggacagcaa ggacagcacc tacagcctca gcagcaccct 660
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243 gggcctga 728

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VERIFICATION SUMMARY

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Input Set : A:\PTO.AMC.txt

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